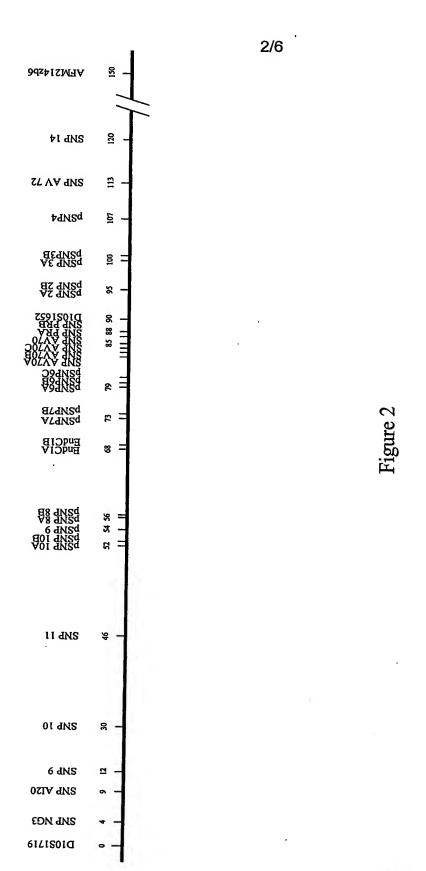


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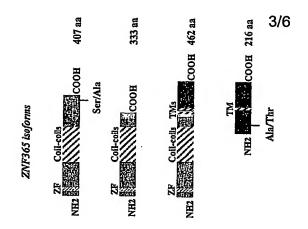
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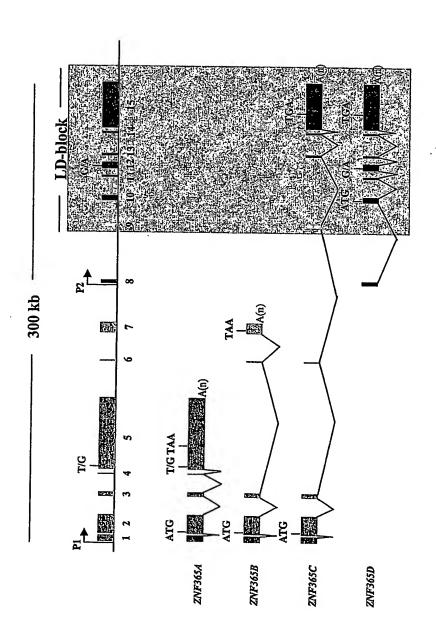


Figure.

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100 100 100	200	300 300 300 54	400 333 400 154	407 462 216
5A MOOKAFEESRYPWOESERNYAVCLDIRCPRCCOHTRRRSIISSIRAHIERSHSYEERFTLIKOSIIRPSIKOTDIVITSSELIKPGKIOSSGNVVKOKPSYVN 5B MOOKAFEESRYPWOESFENVAVCIDIRCPRCCOHTRRRSIISSIRAHIERSHSYEERRIILIKCSIIRPSIKOTDIVITSSELIKPGKIOSSGNVVKOKPSYVN 5C MOOKAREESRYPWOESFENVAVCIDIRCPRCCOHTRERSIISSIRAHERFSHSYEERFTLIKUSIIRPSIIKOTDIVITSSELIKPGKIOSSGNVVKOKPSYVN	LYSISHEHSKORKPERVANAERPVSKVOTVITAMDLHADSIDGIRSGEGILPTSDIKASFRAHVREKENNWEAVDRTLEKKIDK DYSISHEHSKORKPERVARRPVSKVOTYTAMDEHADSIDGIRSGEGIEDTSDIKASEEAHVREKENRWEAVDRUFEKRIDK LYSISHEHSKORKPEEWWERPVSKVOTYTAMDIHADSIDGIRSGEGEDTSDIKASEERHVREKENRWEAVDRUFEKRIDK	FVQLTCKKOBVORRERALNROVDVAVEMITAVIRORLTESRERBITRKBERVATENHEIETAAAEKEVOGKARLODETENLIGERVETAERGIEVYOSOOASGE EVOLTCKKOBVORRERALNROVDVAVEMITAVIRORLTESBELITRKSEERWITENHEIETAABKKEVOGKARLODETENTLORVETAERGIEVYOSOOASGE EVOLTCKKOEVORRERALNROVDVAVEMITAVIRORETESBELITRKSEERVITRINHEIETAAABKKEVOGKARLODETENTLORVETAERGIEVYOSOOASGE EVOLTCKKOEVORRERALNROVDVAVEMITAVIRORETESBELITRKEERVITRINHEIETAAABKKEVOGKARLODETENTLORVETAERGIEVYOSOOASGE SVOLTCKKOEVORRERALNROVDVAVEMITAVIRORETESBEEDINKEEEVATIFNHEIETAAABKKEVOGKARLODETENTLORVETAERGIEVYOSOOASGE NSALGOITITVSRCWNTERNOTDKNPCLHGAYLOLRETVKNKSTHLKKPLMKQA *	VRDISGHVITDISSNRKPKGLSRGHPHSVCNHPDLKSHFHPKGRNHLKKAKODRASMQPAKAIHEQAESSRULCRPPKKGELLGFGRKGNIRPKMAKKKP Transmembrane domains NRDISGHVSWKGAGGEARLVCONDIELEIFGHINHHLSGLKDSHCLVFLQAPPVPWIILASFLWILGNPWTSSTATAGFSGLWVTAPPKGGGTEHHNEKDVI PWKDHLAFQPLHPAERKTQVWRWQSGNSSDLETTSSASPWPTGSNRDVVLNTLAESCCGLSELITAPPYAGVSIQGESGTWVTFPFEGGTEHHNEKDVI * **** *	A TAIVNII CHODFERESVSTSOSRNISLITHGOLONCVTCKATTTETTOLITHETHERVPHCVTCFPWGLDSSS D GHODFERESVSTSOSRNISLITHGOLONCVTCKATTTTOLITHETHETHGVRHCVTCFPWGLDSSS # # #
ZNF365A ZNF365B ZNF365C	ZNF365A ZNF365B ZNF365C	ZNF365A ZNF365B ZNF365C ZNF365D	ZNF365A ZNF365B ZNF365C ZNF365D	ZNF365A ZNF365C ZNF365D

Figure 4

Ī	is in	1					
D10S1640	1055	n.s.	n.s.	n.s.	n.s.		
AFM234wc5	850	n.s.	11.5.	n.s.	n.s.		
L dns	527	n.s.	п.s.	n.s.	n.s.		
I ANS	260	n.s.	n.s.	n.s.	n.s.		
N dns	216	n.s.	n.s.	11.8.	n.s.		
VEMT14zpe	175	0,0213	n.s.	n.s.	n.s.	08	
ti ans	120	n.s.	n.s.	n.S.	n.s.	0,0280	
ZLAY dns	113	n.s.	n.s.	n.S.	n.s.		
AE GN29	100	0,0006 0,0123	0,0004 0,0073	0,0009 0,0145	0,0031 0,0075	39 06	٧
D1021652	90	0,0006	0,0004	0,000	0,0031	05 0,0239 0,0306	Figure 5
0LAA GNS	85	n.s.	n.s.	n.S.	n.s.	0,0205	Ţ
AATE DOLL	83	0,0123	9800.0	960000	1500'0		
11 ANS	46	n.s.	n.s.	n.s.			
01 ANS	30	n.s.	n.s.	n.s.	n.s.		
6 dns	12	n.s.	n.s.	n.s.	n.s.		
SNP NG3	4	n.s.	n.s.	n.s.	n.s.	· · · · · · · · · · · · · · · · · · ·	
61/1S01 <b>Q</b>	1	n.S.	n.s.	n.s.	n.s.		
Marcatore	Posizione (Kb)	val. p <sup>a</sup>	val. p empirico <sup>b</sup>	val. p <sup>c</sup>	val. p empirico <sup>d</sup>	val. p empirico <sup>e</sup> (analisi aplotipo)	

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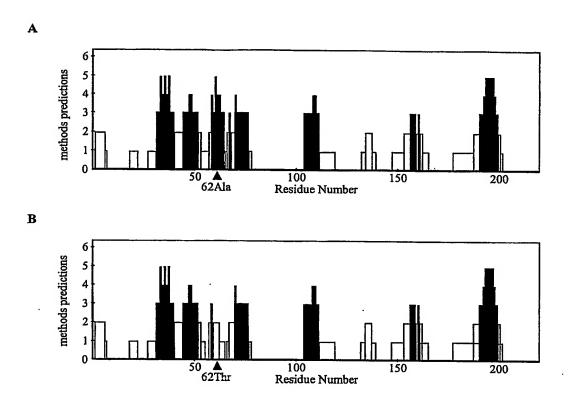


Figure 6

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